RAW SEQUENCE LISTING PATENT APPLICATION US/08/948,124

DATE: 07/10/98 TIME: 15:33:04

	INPUT SET: S27370.raw
	This Raw Listing contains the General Information Section and up to the first pages. SEQUENCE LISTING (1) General Information:
1 2	SEQUENCE LISTING
3	(1) General Information:
4 5 6 7 8	(i) APPLICANT: Reinherz, Ellis Clayton, Linda Ocain, Timothy D. Patch, Raymond J.
9 10 11 12	(ii) TITLE OF INVENTION: THYMOCYTE CASPASE ACTIVITY AND NEGATIVE SELECTION
13 14	(iii) NUMBER OF SEQUENCES: 9
15 16 17	(iv) CORRESPONDENCE ADDRESS:(A) ADDRESSEE: HAMILTON, BROOK, SMITH & REYNOLDS, P.C.(B) STREET: Two Militia Drive
18 19 20	(C) CITY: Lexington (D) STATE: MA (E) COUNTRY: USA
21 22	(F) ZIP: 02173
23 24	(V) COMPUTER READABLE FORM: (A) MEDIUM TYPE: Diskette
25 26 27	(B) COMPUTER: IBM Compatible (C) OPERATING SYSTEM: Windows 95 (D) SOFTWARE: FastSEQ for Windows Version 2.0b
28 29	(vi) CURRENT APPLICATION DATA:
30 31 32	(A) APPLICATION NUMBER: 08/948,124 (B) FILING DATE: 09-OCT-1997 (C) CLASSIFICATION:
33 34 35	(Vii) PRIOR APPLICATION DATA: (A) APPLICATION NUMBER: 08/802,474
36 37 38 39	(B) FILING DATE: 18-FEB-1997
40 41 42	<pre>(viii) ATTORNEY/AGENT INFORMATION: (A) NAME: Carroll, Alice O (B) REGISTRATION NUMBER: 33,542</pre>
43 44	(C) REFERENCE/DOCKET NUMBER: DFCI-522A
45 46	(ix) TELECOMMUNICATION INFORMATION: (A) TELEPHONE: 781-861-6240

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47
              (B) TELEFAX: 781-861-9540
              (C) TELEX:
48
49
50
               (2) INFORMATION FOR SEQ ID NO:1:
51
52
53
            (i) SEQUENCE CHARACTERISTICS:
              (A) LENGTH: 5 amino acids
55
              (B) TYPE: amino acid
              (C) STRANDEDNESS: single
56
57
              (D) TOPOLOGY: linear
58
59
            (ii) MOLECULE TYPE: peptide
60
            (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:
61
62
63
      Gln Ala Cys Xaa Gly
64
65
66
               (2) INFORMATION FOR SEQ ID NO:2:
67
68
            (i) SEQUENCE CHARACTERISTICS:
              (A) LENGTH: 4 amino acids
69
70
              (B) TYPE: amino acid
              (C) STRANDEDNESS: single
71
72
              (D) TOPOLOGY: linear
73
74
            (ii) MOLECULE TYPE: peptide
75
76
            (xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:
77
78
     Tyr Val Ala Asp
79
80
81
               (2) INFORMATION FOR SEQ ID NO:3:
82
83
            (i) SEQUENCE CHARACTERISTICS:
84
              (A) LENGTH: 4 amino acids
85
              (B) TYPE: amino acid
              (C) STRANDEDNESS: single
86
              (D) TOPOLOGY: linear
87
88
89
            (ii) MOLECULE TYPE: peptide
90
            (xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:
91
92
93
     Asp Glu Val Asp
94
95
96
               (2) INFORMATION FOR SEQ ID NO:4:
97
98
            (i) SEQUENCE CHARACTERISTICS:
              (A) LENGTH: 9 amino acids
```

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	nw o	1 511. 52/5/0.14#
100	(B) TYPE: amino acid	
101	(C) STRANDEDNESS: single	
102	(D) TOPOLOGY: linear	
103	• •	
104	(ii) MOLECULE TYPE: peptide	
105	(11, 110120011 11110	
106	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:4:	
	(XI) SEQUENCE DESCRIPTION: SEQ ID NO.4.	
107	mi ali mi dli ani min min ali fan	
108	Phe Ala Pro Gly Asn Tyr Pro Ala Leu	
109	1 5	
110		
111	(2) INFORMATION FOR SEQ ID NO:5:	
112		
113	(i) SEQUENCE CHARACTERISTICS:	
114	(A) LENGTH: 100 base pairs	
115	(B) TYPE: nucleic acid	
116	(C) STRANDEDNESS: single	
117	(D) TOPOLOGY: linear	
	(D) TOPOLOGI: Timedi	
118		
119	(') CROUNTED PROCEETINGS OF THE NO. F.	
120	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:5:	
121		
122	CTCGAGACCT GTGTGGATTA AAAACCTCTC TGATTCTGGT TTGCTTTTCT GT	
123	AGGGGTGAGA GGAGACTCCG TGACCCAGAC AGAAGGCCTG	100
124		
125	(2) INFORMATION FOR SEQ ID NO:6:	
126		
127	(i) SEQUENCE CHARACTERISTICS:	
128	(A) LENGTH: 69 base pairs	
129	(B) TYPE: nucleic acid	
130	(C) STRANDEDNESS: single	
131	(D) TOPOLOGY: linear	
	(D) TOPOLOGI: IIMeal	
132		
133		
134	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:6:	
135		
136	GCGGCCGCTT GGGCCCAAGA AACTGTCATC AAACGTACGT GCTATAACCT TC	
137	ACCTGCTCC	69
138		
139	(2) INFORMATION FOR SEQ ID NO:7:	
140		
141	(i) SEQUENCE CHARACTERISTICS:	
142	(A) LENGTH: 86 base pairs	
143	(B) TYPE: nucleic acid	
144	(C) STRANDEDNESS: single	
144	(D) TOPOLOGY: linear	
	(D) TOPOLOGI. IIMEGI	
146		
147	A LA ARAMBUAR BURARTER AND TO MA T	
148	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:7:	
149		
150	ATCGATTCCC CTTCTTGTCT GTATTTACAT CTTTTGTTTT CTTCTCTTCT	
151	GATTCTGGGG TTGTCCAGTC TCCAAG	86
152		

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	1141 U1 3E1. 32/3/V.14W
153	(2) INFORMATION FOR SEQ ID NO:8:
154	
155	(i) SEQUENCE CHARACTERISTICS:
156	(A) LENGTH: 56 base pairs
157	(B) TYPE: nucleic acid
158	(C) STRANDEDNESS: single
159	(D) TOPOLOGY: linear
160	
161	
162	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:8:
163	
164	GCGGCCGCAC GTGGGGCCCC AGCTCACCTA AAACCGTGAG CCTGGTGCCG GGACCG 56
165	
166	(2) INFORMATION FOR SEQ ID NO:9:
167	
168	(i) SEQUENCE CHARACTERISTICS:
169	(A) LENGTH: 8 amino acids
170	(B) TYPE: amino acid
171	(C) STRANDEDNESS: single
172	(D) TOPOLOGY: linear
173	
174	(ii) MOLECULE TYPE: peptide
175	
176	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:9:
177	
178	Arg Gly Tyr Val Tyr Gln Gly Leu
179	1 5

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SEQUENCE VERIFICATION REPORT PATENT APPLICATION US/08/948,124

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